

FIGURE 1

1 gtgaaacacc ctcggctggg aagtcagttc gttctctcct ctcctctctt cttgtttgaa
61 catggtgcgg actaaagcag acagtgttcc aggcacttac agaaaagtgg tggctgctcg
121 agcccccaga aagggtgcttg gttcttccac ctctgccact aattcgacat cagtttcatc
181 gagggaaagct gaaaataaat atgcaggagg gaaccccggtt tgcgtgcgcc caactcccaa
241 gtggcaaaaa ggaattggag aattcttttag gttgtccctt aaagattctg aaaaagagaa
301 tcagattcct gaagaggcag gaagcagtgg cttagaaaa gcaaagagaa aagcatgtcc
361 tttgcaacct gatcacacaa atgatgaaaa agaatagaac tttctcattc atctttgaat
421 aacgtctcct tggttaccct ggtattctag aatgtaaatt tacataaatg tggttggcc
481 aattagctt gttgaacagg catttaatta aaaaatttag gtttaaattt agatgttcaa
541 aagttagttgt gaaatttgag aatttgtaaag actaattatg gtaacttagc ttagtattca
601 atataatgca ttgtttgggtt tcttttacca aattaagtgt cttagtcttg ctaaaatcaa
661 gtcattgcat tgggttctaa ttacaagtat gttgtattt agatttgctt agattgttgt
721 actgctgcca tttttattgg tgggttattttt gttttgatta ttggaaatggt gccatattgt cactccttct
781 acttgcttta aaaagcagag tttagattttt gcacattaaa aaattcaaga ttaatt

FIGURE 2A

MVRTKADSSVPGTYRKVVAAARAPRKVILGSSTSATNSTSVSSRK
AENKYAGGNPVCVRPTPKWQKGIGEFFRLSPKDSEKENQIPE
EAGSSGLGKAKRKACPLQPDHTNDEKE

p15PAF
XPG (981-1009) LKQLDAQQQTQLRIDSEFFRLAQQEKEDAKR
FEN-1 (328-355) RQGSTQGRILDDEFEKVTGSL
P21 (139-160) KRRQTSMTDIFYHSKRRLIES
MCMT (162-174) TRQTTTSHFAKG.

FIGURE 2B

FIGURE 3

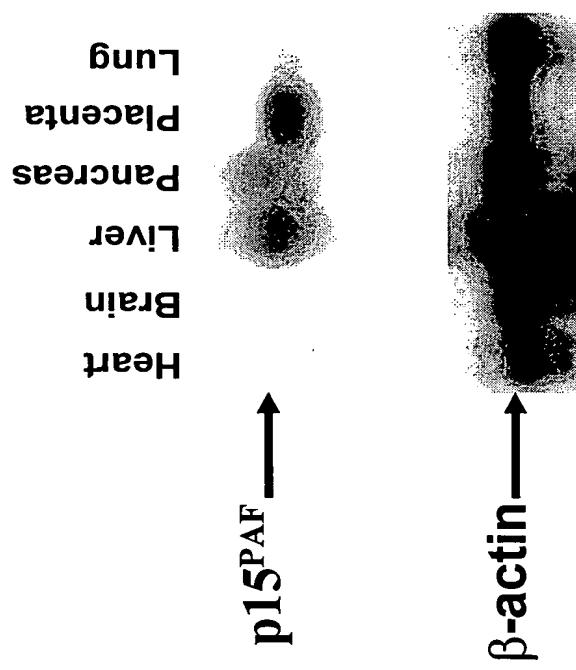


FIGURE 4

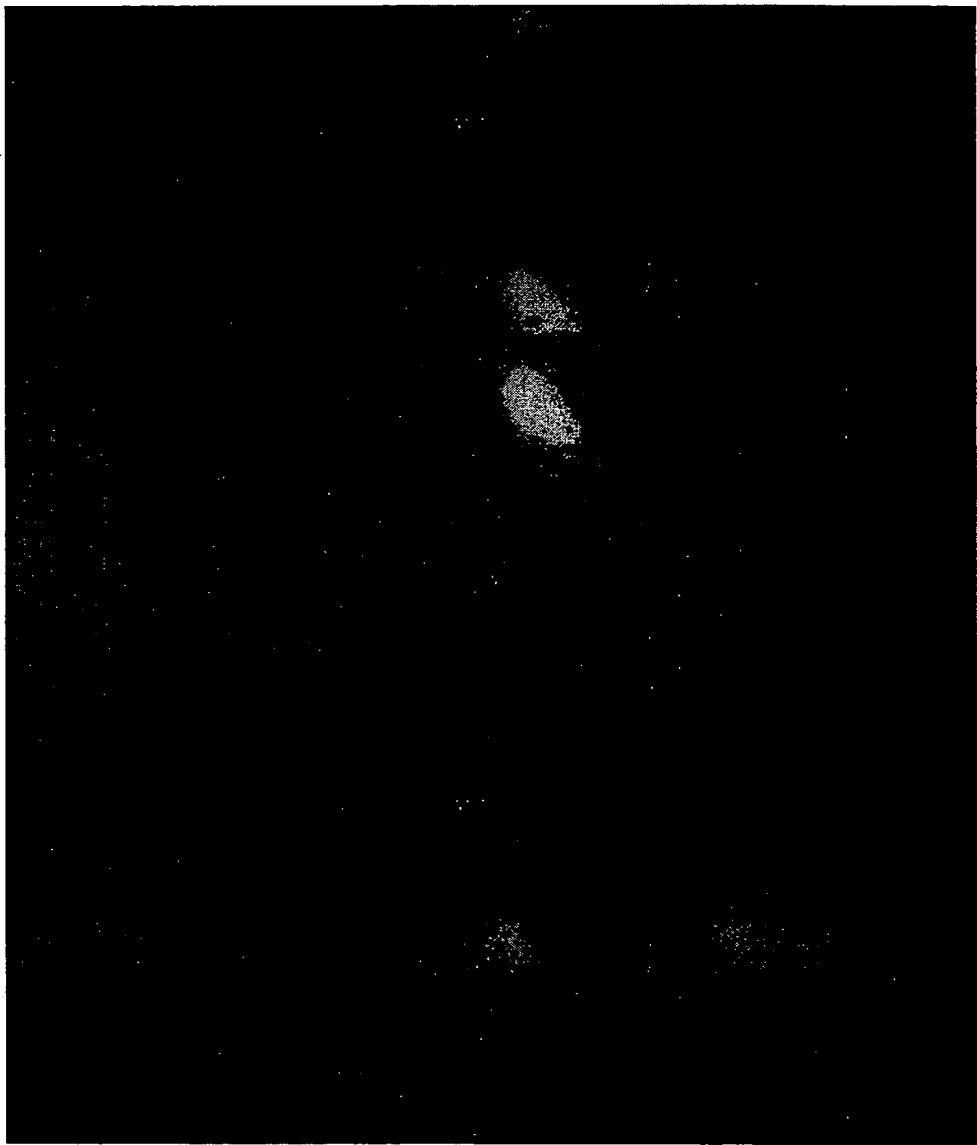


FIGURE 5

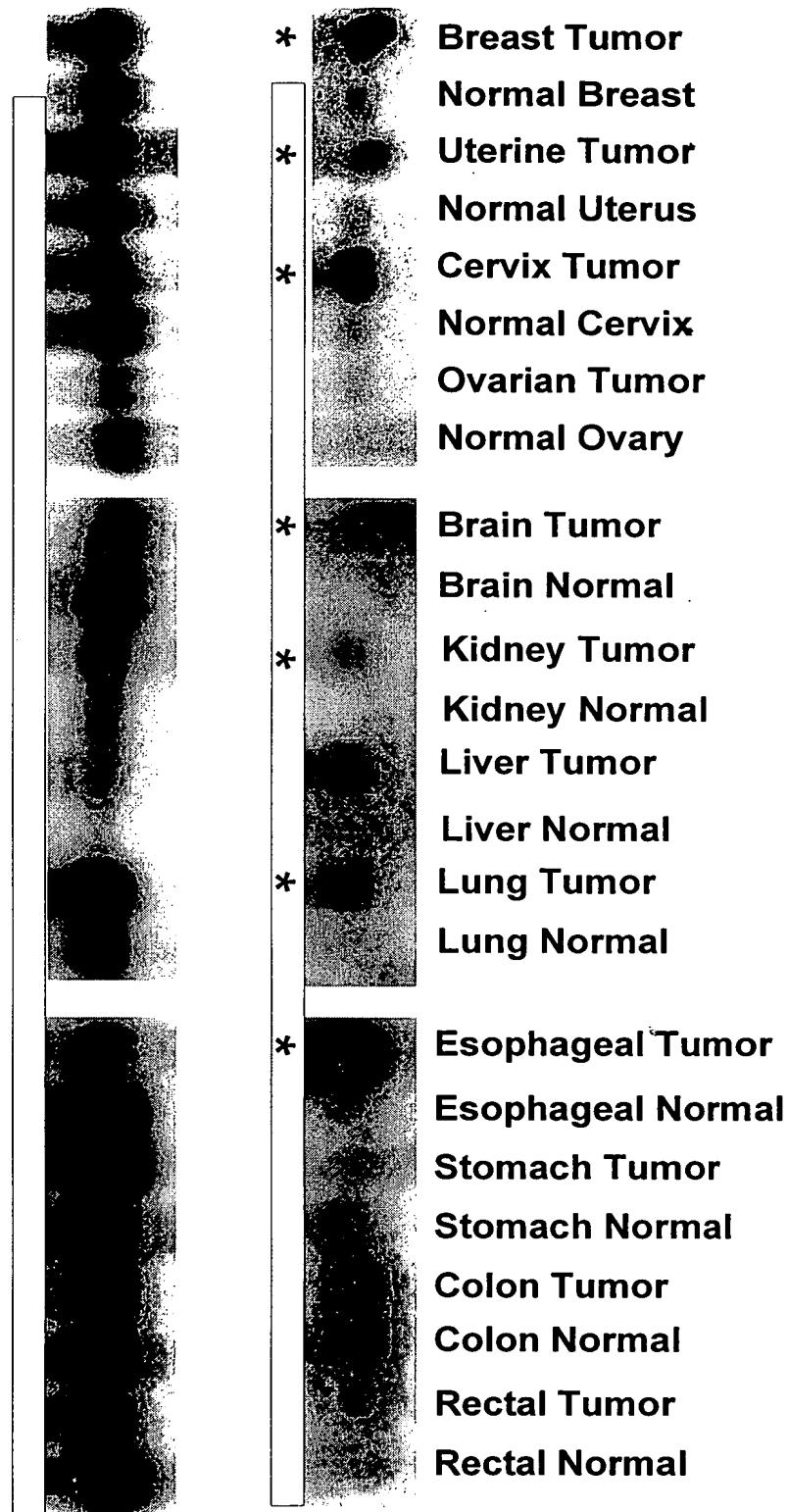


FIGURE 6

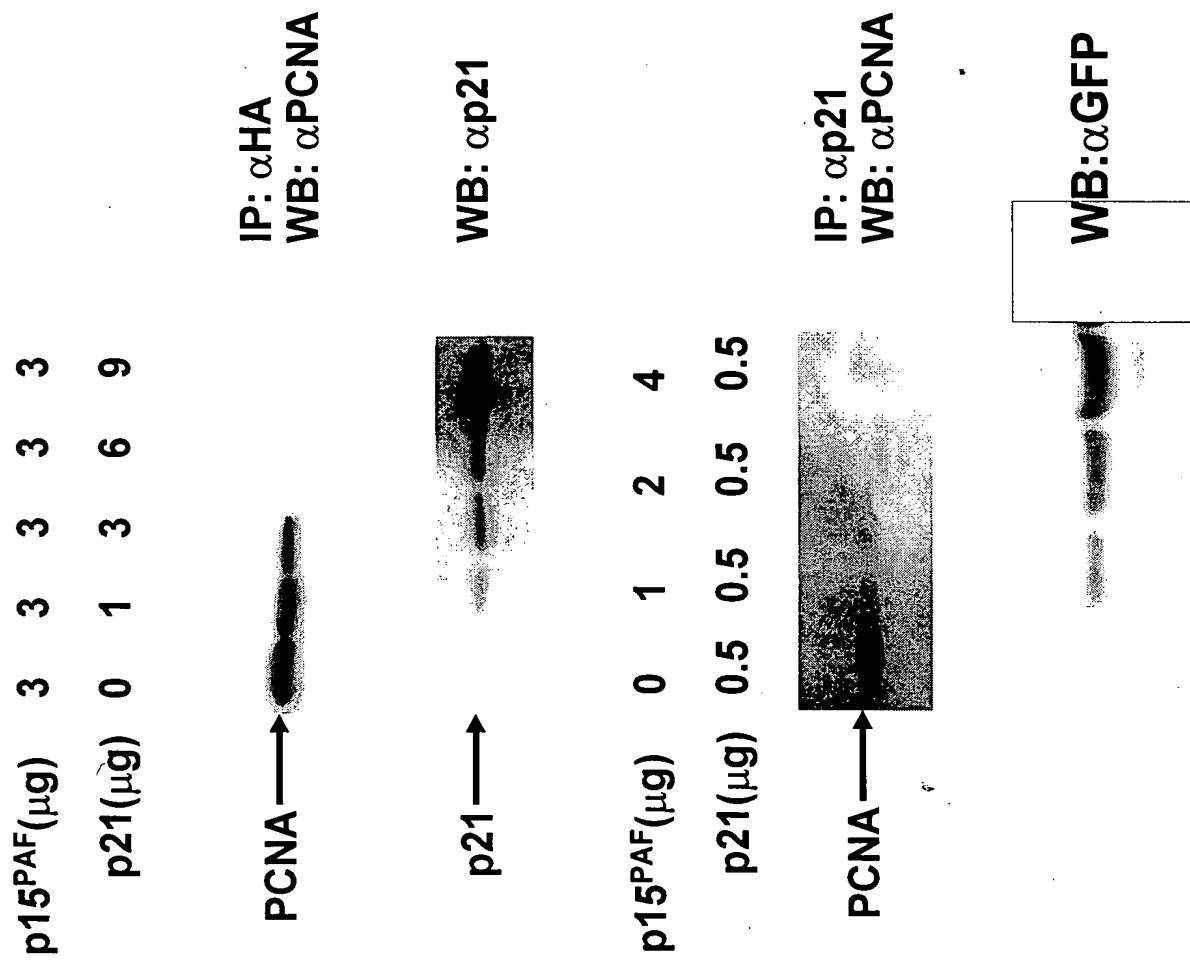


FIGURE 7

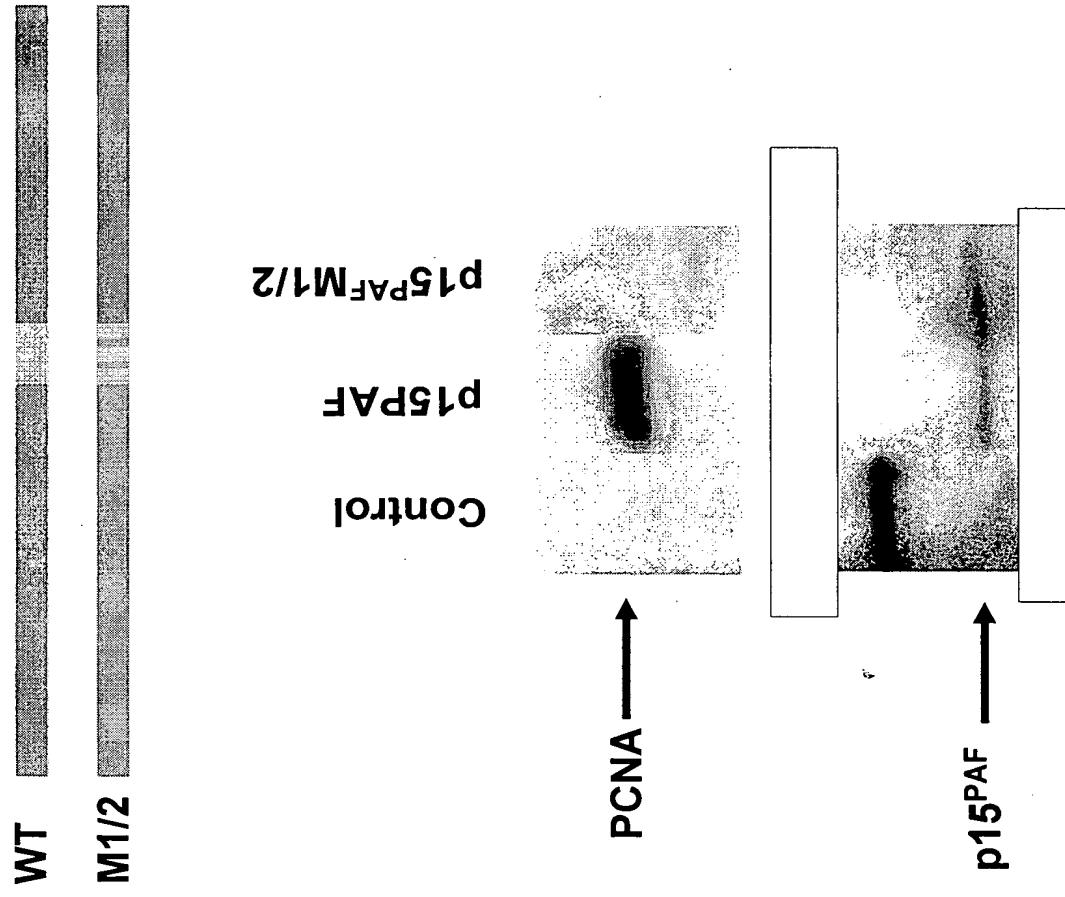


FIGURE 8

R0101 binds to PCNA, CDK2, CDK3

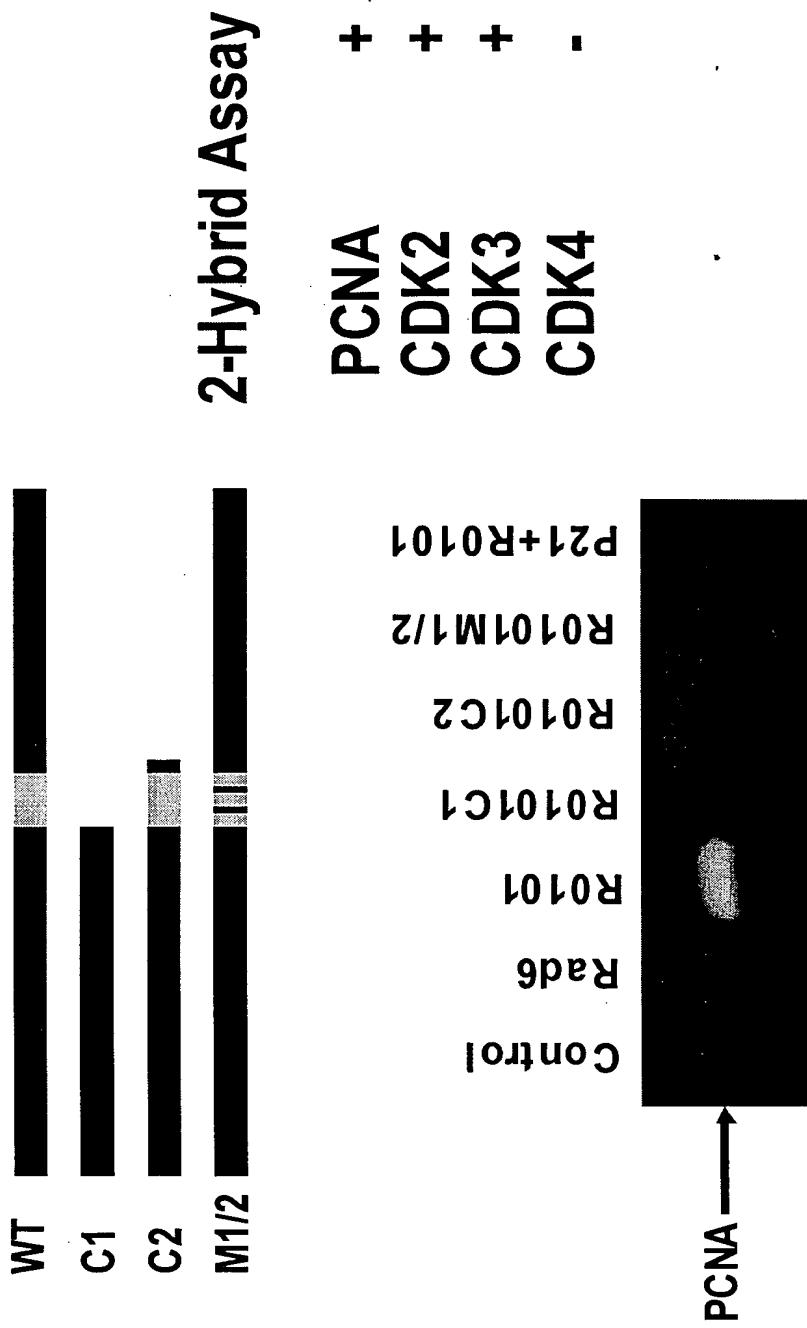


FIGURE 9

